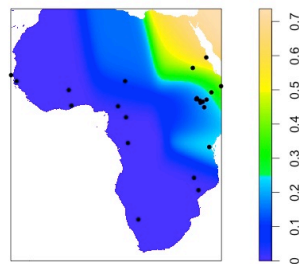


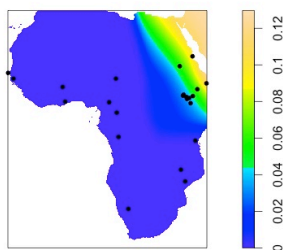
Supplementary Figure 3

Contour maps showing the distribution of the 4 most frequent individual LP functional enhancer alleles in comparison with the distribution of LP frequency calculated from all five functional alleles together. *-13910*T* is too rare to make a meaningful map. Note that for *-13907*G* particularly, the absence of data points in the Middle East in this study slightly distorts the contour map. Fuller maps will be shown in our next publication: Liebert et al in preparation.

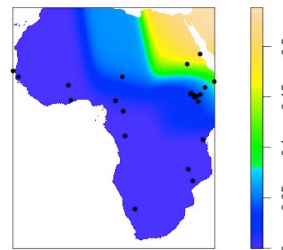
LP calculated from 5 alleles



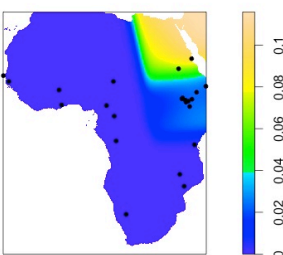
Allele frequency *-13907*G*



Allele frequency *-13915*G*



Allele frequency *-14009*G*



Allele frequency *-14010*C*

